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Range: from to ☐ Reverse complemented strand Features:
☐ 1: [NM_080751](#). Reports Homo sapiens tran...[gi:94536851]

[Links](#)
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LOCUS NM_080751 3169 bp mRNA linear PRI [14-MAY-2006](#)

DEFINITION Homo sapiens transmembrane channel-like 2 (TMC2), mRNA.

ACCESSION NM_080751

VERSION NM_080751.2 GI:94536851

KEYWORDS .

SOURCE Homo sapiens (human)

ORGANISM Homo sapiens

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Primates; Haplorrhini; Catarrhini; Hominidae; Homo.

REFERENCE 1 (bases 1 to 3169)

AUTHORS Kimura,K., Wakamatsu,A., Suzuki,Y., Ota,T., Nishikawa,T., Yamashita,R., Yamamoto,J., Sekine,M., Tsuritani,K., Wakaguri,H., Ishii,S., Sugiyama,T., Saito,K., Isono,Y., Irie,R., Kushida,N., Yoneyama,T., Otsuka,R., Kanda,K., Yokoi,T., Kondo,H., Wagatsuma,M., Murakawa,K., Ishida,S., Ishibashi,T., Takahashi-Fujii,A., Tanase,T., Nagai,K., Kikuchi,H., Nakai,K., Isogai,T. and Sugano,S.

TITLE Diversification of transcriptional modulation: large-scale identification and characterization of putative alternative promoters of human genes

JOURNAL Genome Res. 16 (1), 55-65 (2006)

PUBMED [16344560](#)

REFERENCE 2 (bases 1 to 3169)

AUTHORS Kurima,K., Yang,Y., Sorber,K. and Griffith,A.J.

TITLE Characterization of the transmembrane channel-like (TMC) gene family: functional clues from hearing loss and epidermodysplasia verruciformis

JOURNAL Genomics 82 (3), 300-308 (2003)

PUBMED [12906855](#)

REFERENCE 3 (bases 1 to 3169)

AUTHORS Keresztes,G., Mutai,H. and Heller,S.

TITLE TMC and EVER genes belong to a larger novel family, the TMC gene family encoding transmembrane proteins

JOURNAL (er) BMC Genomics 4 (1), 24 (2003)

PUBMED [12812529](#)

REFERENCE 4 (bases 1 to 3169)

AUTHORS Kurima,K., Peters,L.M., Yang,Y., Riazuddin,S., Ahmed,Z.M., Naz,S., Arnaud,D., Drury,S., Mo,J., Makishima,T., Ghosh,M., Menon,P.S., Deshmukh,D., Oddoux,C., Ostrer,H., Khan,S., Riazuddin,S., Deininger,P.L., Hampton,L.L., Sullivan,S.L., Battey,J.F. Jr., Keats,B.J., Wilcox,E.R., Friedman,T.B. and Griffith,A.J.

TITLE Dominant and recessive deafness caused by mutations of a novel gene, TMCL1, required for cochlear hair-cell function

JOURNAL Nat. Genet. 30 (3), 277-284 (2002)

PUBMED [11850618](#)

COMMENT VALIDATED REFSEQ: This record has undergone preliminary review of the sequence, but has not yet been subject to final review. The reference sequence was derived from [AF417580.2](#), [DA769512.1](#) and [AL049712.12](#).

On May 4, 2006 this sequence version replaced gi:[20304092](#).

Summary: This gene is considered a member of a gene family predicted to encode transmembrane proteins. The specific function of this gene is unknown; however, expression in the inner ear suggests that it may be crucial for normal auditory function. Mutations in this gene may underlie hereditary disorders of balance and hearing.

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Range: from to ☐ Reverse complemented strand Features:

☐ 1: [AF417580](#). Reports Homo sapiens tran...[gi:28642834]

[Links](#)

[Comment](#) [Features](#) [Sequence](#)

LOCUS AF417580 3169 bp mRNA linear PRI 05-MAR-2003
 DEFINITION Homo sapiens transmembrane channel-like protein 2 (TMC2) mRNA, complete cds.
 ACCESSION AF417580
 VERSION AF417580.2 GI:28642834
 KEYWORDS .
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 REFERENCE 1 (bases 1 to 3169)
 AUTHORS Kurima,K., Peters,L.M., Yang,Y., Riazuddin,S., Ahmed,Z.M., Naz,S., Arnaud,D., Drury,S., Mo,J., Makishima,T., Ghosh,M., Menon,P.S.N., Deshmukh,D., Oddoux,C., Ostrer,H., Khan,S., Raizuddin,S., Deininger,P.L., Hampton,L.L., Sullivan,S.L., Battey,J.F., Keats,B.J.B., Wilcox,E.R., Friedman,T.B. and Griffith,A.J.
 TITLE Dominant and recessive deafness caused by mutations of a novel gene, TMC1, required for cochlear hair-cell function
 JOURNAL Nat. Genet. 30 (3), 277-284 (2002)
 PUBMED [11850618](#)
 REFERENCE 2 (bases 1 to 3169)
 AUTHORS Kurima,K., Griffith,A.J. and Friedman,T.B.
 TITLE Direct Submission
 JOURNAL Submitted (10-SEP-2001) NIDCD, NIH, 5 Research Court, #2A02, Rockville, MD 20850, USA
 REFERENCE 3 (bases 1 to 3169)
 AUTHORS Kurima,K., Griffith,A.J. and Friedman,T.B.
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 JOURNAL Submitted (03-MAR-2003) NIDCD, NIH, 5 Research Court, #2A02, Rockville, MD 20850, USA
 REMARK Sequence update by submitter
 COMMENT On Mar 3, 2003 this sequence version replaced gi:19223982.
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Apr 11 2006 19:57:30

AF417580
LOCUS AF417580 3169 bp mRNA linear PRI 05-MAR-2003
DEFINITION Homo sapiens transmembrane channel-like protein 2 (TMC2) mRNA, complete cds.
ACCESSION AF417580
VERSION AF417580.2 GI:28642834
KEYWORDS .
SOURCE Homo sapiens (human)
ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae; Homo.
REFERENCE 1 (bases 1 to 3169)
AUTHORS Kurima,K., Peters,L.M., Yang,Y., Riazuddin,S., Ahmed,Z.M., Naz,S., Arnaud,D., Drury,S., Mo,J., Makishima,T., Ghosh,M., Menon,P.S.N., Deshmukh,D., Oddoux,C., Ostrer,H., Khan,S., Raizuddin,S., Deininger,P.L., Hampton,L.L., Sullivan,S.L., Battey,J.F., Keats,B.J.B., Wilcox,E.R., Friedman,T.B. and Griffith,A.J.
TITLE Dominant and recessive deafness caused by mutations of a novel gene, TMC1, required for cochlear hair-cell function
JOURNAL Nat. Genet. 30 (3), 277-284 (2002)
PUBMED 11850618
REFERENCE 2 (bases 1 to 3169)
AUTHORS Kurima,K., Griffith,A.J. and Friedman,T.B.
TITLE Direct Submission
JOURNAL Submitted (10-SEP-2001) NIDCD, NIH, 5 Research Court, #2A02, Rockville, MD 20850, USA
REFERENCE 3 (bases 1 to 3169)
AUTHORS Kurima,K., Griffith,A.J. and Friedman,T.B.
TITLE Direct Submission
JOURNAL Submitted (03-MAR-2003) NIDCD, NIH, 5 Research Court, #2A02, Rockville, MD 20850, USA
REMARK Sequence update by submitter
COMMENT On Mar 3, 2003 this sequence version replaced gi:19223982.
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ORIGIN

Query Match 100.0%; Score 3169; DB 8; Length 3169;
Best Local Similarity 100.0%; Pred. No. 0;
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Qy	61	GTGAAAGGGCGGGTGAAGAGCGGCTCTCCACACACAGGTGACAGGCTGGGAAGGAGATCC	120
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Qy	241	CGCAGGAGACACAGAGAAGAGCTGGGGGAGCAGGAGCGGGGCGAGGCAGAGAGGACCTGC	300
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Qy	301	GAGGGCAGGAGAAAGCGCGACGAGAGGGCCTCCTTCCAGGAGCGGACAGCAGCCCCAAAG	360
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Qy	481	ATCCTGGAGCAGGTGGAAGAAAAAAGAAGCTCATTGCCACCATGCGGAGCAAGCCCTGG	540
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Qy	541	CCCATGGCGAAGAAGCTGACAGAGCTCAGGGAGGCCAGGAATTTGTGGAGAAGTATGAA	600
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Qy	961	TACTCTGCACTCTTCTATGGCTACTACAACAACCAGAGGACCATCGGGTGGCTGAGGTAC	1020
Db	961	TACTCTGCACTCTTCTATGGCTACTACAACAACCAGAGGACCATCGGGTGGCTGAGGTAC	1020
Qy	1021	CGGCTGCCTATGGCTTACTTTATGGTGGGGGTACGCGTGTTTCGGCTACAGCCTGATTATT	1080
Db	1021	CGGCTGCCTATGGCTTACTTTATGGTGGGGGTACGCGTGTTTCGGCTACAGCCTGATTATT	1080
Qy	1081	GTCATTTCGATCGATGGCCAGCAATACCCAAGGAAGCACAGGCGAAGGGGAGAGTGACAAC	1140
Db	1081	GTCATTTCGATCGATGGCCAGCAATACCCAAGGAAGCACAGGCGAAGGGGAGAGTGACAAC	1140
Qy	1141	TTCACATTTCAGCTTCAAGATGTTTACCAGCTGGGACTACCTGATCGGGAATTTCAGAGACA	1200
Db	1141	TTCACATTTCAGCTTCAAGATGTTTACCAGCTGGGACTACCTGATCGGGAATTTCAGAGACA	1200
Qy	1201	GCTGATAACAAATATGCATCCATCACCACCAGCTTCAAGGAATCAATAGTGGATGAACAA	1260
Db	1201	GCTGATAACAAATATGCATCCATCACCACCAGCTTCAAGGAATCAATAGTGGATGAACAA	1260
Qy	1261	GAGAGTAACAAAGAAGAAAATATCCATCTGACAAGATTTCTTCGTGTCCTGGCCAACTTT	1320
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Qy	1381	CAGCAATTCTCCAAAATGCAGAATGTCAGCTGGTATGAAAGGAATGAGGTAGAGATCGTG	1440
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Qy	1501	TACCACCCACGCACTGGACTGAAGTGGCAGCTGGGACGCATCTTTGCACTCTTCCTGGGG	1560
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Qy	1561	AACCTCTACACATTTCTCTTGGCCCTGATGGATGACGTCCACCTCAAGCTTGCTAATGAA	1620
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Qy	1801	ATCACCATCCTGCTGGGGGACTTCCTACGGGCTTGTTTTGTGCGGTTTCATGAACACTGC	1860
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Qy	1861	TGGTGCTGGGACTTGGAGGCTGGATTTCTTCATATGCTGAGTTTGATATTAGTGGAAT	1920
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Qy	1921	GTGCTGGGTTTGATCTTCAACCAAGGAATGATCTGGATGGGCTCCTTCTATGCTCCAGGC	1980
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Qy	1981	CTGGTGGGCATTAATGTGCTGCGCCTGCTGACCTCCATGTACTTCCAGTGCTGGGCGGTG	2040
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Qy	3121	AACCAGGAGGGGGATGGAGAAGGGCCTACATTTCTCAATCCAGAGGAAG	3169
Db	3121	AACCAGGAGGGGGATGGAGAAGGGCCTACATTTCTCAATCCAGAGGAAG	3169



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bacteria: 0

mRNA: 0

RefSeq: 0



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